

# United States Patent and Trademark Office



UNITED STATES DEPARTMENT OF COMMERCE United States Patent and Trademark Office Address: COMMISSIONER FOR PATENTS P.O. Box 1450 Alexandria, Virginia 22313-1450 www.uspto.gov

APPLICATION NO.	F	ILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.	
09/910,383	910,383 07/20/2001		Girish N. Nallur	13172.0007U1	2312	
23859	7590	06/15/2006		EXAMINER		
		NBERG, P.C.	CALAMITA, HEATHER			
SUITE 1000 999 PEACHTREE STREET ATLANTA, GA 30309-3915				ART UNIT	PAPER NUMBER	
				1637		
				DATE MAILED: 06/15/2006		

Please find below and/or attached an Office communication concerning this application or proceeding.

		A multipation No.	I Applicant (a)			
Office Action Summary		Application No.	Applicant(s)			
		09/910,383	NALLUR ET AL.			
		Examiner	Art Unit			
		Heather G. Calamita, Ph.D.	1637			
Period fo	The MAILING DATE of this communication app or Reply	ears on the cover sheet with the d	orrespondence address			
WHIC - External after - If NO - Failu Any r	ORTENED STATUTORY PERIOD FOR REPLY CHEVER IS LONGER, FROM THE MAILING DATE in may be available under the provisions of 37 CFR 1.13 SIX (6) MONTHS from the mailing date of this communication. Operiod for reply is specified above, the maximum statutory period we are to reply within the set or extended period for reply will, by statute, reply received by the Office later than three months after the mailing ed patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be tiruit apply and will expire SIX (6) MONTHS from cause the application to become ABANDONE	N. mely filed the mailing date of this communication. ED (35 U.S.C. § 133).			
Status						
1)⊠	Responsive to communication(s) filed on 22 M	arch 2006.				
2a) <u></u> □	This action is <b>FINAL</b> . 2b)⊠ This	action is non-final.				
3) 🗌	Since this application is in condition for allowance except for formal matters, prosecution as to the merits is					
	closed in accordance with the practice under E	x parte Quayle, 1935 C.D. 11, 4	53 O.G. 213.			
Dispositi	ion of Claims					
5)□ 6)⊠ 7)□	Claim(s) 1-62,68-75 and 82-87 is/are pending if 4a) Of the above claim(s) is/are withdraw Claim(s) is/are allowed.  Claim(s) 1-62,68-75 and 82-87 is/are rejected.  Claim(s) is/are objected to.  Claim(s) are subject to restriction and/or	vn from consideration.				
Applicati	ion Papers					
• —	The specification is objected to by the Examine The drawing(s) filed on is/are: a) access Applicant may not request that any objection to the Replacement drawing sheet(s) including the correction	epted or b) objected to by the drawing(s) be held in abeyance. Se	e 37 CFR 1.85(a).			
11)	The oath or declaration is objected to by the Ex	* * * * * * * * * * * * * * * * * * * *				
Priority u	ınder 35 U.S.C. § 119					
12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).  a) All b) Some * c) None of:  1. Certified copies of the priority documents have been received.  2. Certified copies of the priority documents have been received in Application No  3. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).  * See the attached detailed Office action for a list of the certified copies not received.						
2)  Notic Notic  Notic	t(s) te of References Cited (PTO-892) te of Draftsperson's Patent Drawing Review (PTO-948) mation Disclosure Statement(s) (PTO-1449 or PTO/SB/08) r No(s)/Mail Date	4) Interview Summary Paper No(s)/Mail D 5) Notice of Informal F 6) Other:				

#### **DETAILED ACTION**

## Continued Examination Under 37 CFR 1.114

1. A request for continued examination under 37 CFR 1.114, including the fee set forth in 37 CFR 1.17(e), was filed in this application after final rejection. Since this application is eligible for continued examination under 37 CFR 1.114, and the fee set forth in 37 CFR 1.17(e) has been timely paid, the finality of the previous Office action has been withdrawn pursuant to 37 CFR 1.114. Applicant's submission filed on March 22, 2006, has been entered.

## Status of Application, Amendments, and/or Claims

2. Claims 1-62, 68-76 and 82-87 are pending and under examination. All arguments have been fully considered and thoroughly reviewed, but are deemed not persuasive for the reasons that follow. Any objections and rejections not reiterated below are hereby withdrawn.

## Claim Rejections - 35 USC § 103

- 3. The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:
  - (a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.

Claims 1-15, 18-29, 31-47, 53-58, 61, 62, 68, 70-73 and 82-87 are rejected under 35 U.S.C. 103(a) as being unpatentable over Lizardi et al. (USPN 6,316,229 B1) in view of Schweitzer et al. (PNAS 2000).

Lizardi et al. teach (claims 1, 53, 56, 62, 68, 70-73 and 87) a method of amplifying messenger RNA, the method comprising (see entire document, specifically col. 42 line 13).

Art Unit: 1637

- (a) mixing one or more RT primers with a nucleic acid sample and reverse transcribing to produce cDNA strands each comprising one of the RT primers, wherein each RT primer comprises a reverse transcription primer portion (see col. 77 line 2, where Lizardi et al. indicate RNA can be used with the methodologies disclosed).
- (b) mixing the cDNA strands with a set of capture probes under conditions that promote hybridization of the cDNA strands to the capture probes (see col. 42 lines 27-52)
- (c) mixing one or more rolling circle replication primers with the cDNA strands under conditions that promote association of the cDNA strands with the rolling circle replication primers, wherein the rolling circle replication primers (see col. 42 lines 27-52),
- (d) mixing one or more amplification target circles with the rolling circle replication primers under conditions that promote association of the rolling circle replication primers with the amplification target circles (see col. 42 lines 53-67),
- (e) incubating the amplification target circles under conditions that promote replication of the amplification target circles(see col. 42 lines 65-67, col. 43 lines 1-4), wherein replication of the amplification target circles results in the formation of tandem sequence DNA (see col. 43 lines 5-20).

With regard to claim 3, Lizardi et al. teach the reverse transcription primer portion of each RT primer comprises poly T (see col. 77 line 65).

With regard to claim 4, Lizardi et al. teach the capture probes are immobilized on a substrate (see col. 42 lines 29-30).

With regard to claim 5, Lizardi et al. teach the capture probes are in an array (see col. 42 line 8).

With regard to claim 6, Lizardi et al. teach the capture probes are immobilized via a capture tag coupled to the capture probes (see example 5 col. 78 steps 1 and 2).

With regard to claim 7, Lizardi et al. teach each capture probe comprises a sequence matching all or a portion of the sequence of messenger RNA molecules of interest (see col. 77 line 65, col. 78 line 4).

Art Unit: 1637

With regard to claims 8-12, Lizardi et al. teach the set of capture probes collectively comprise sequence matching all or a portion of the sequence of a plurality of different messenger RNA molecules associated with a disease from a source of interest (see col. 51 lines 63-67, col. 52 lines 1-40).

With regard to claim 13, Lizardi et al. teach the ends of the capture probes are extendable when a cDNA strand is hybridized to the capture probe (see col. 42 line 67).

With regard to claim 14, Lizardi et al. teach the ends of the capture probes are designed to be extendable only when a cDNA strand corresponding to a particular form of a messenger RNA of interest is hybridized to the capture probe (see col. 42 line 67).

With regard to claim 15, Lizardi et al. teach the ends of the capture probes are not extendable by polymerase (see col. 47 lines 62-64).

With regard to claim 18, Lizardi et al. teach further comprising, simultaneous with, or following, step (d), mixing a secondary DNA strand displacement primer with the amplification target circles and incubating under conditions that promote hybridization between the tandem sequence DNA and the secondary DNA strand displacement primer and replication of the tandem sequence DNA, wherein replication of the tandem sequence DNA results in the formation of secondary tandem sequence DNA (see col. 55 lines 40-54).

With regard to claim 19, Lizardi et al. teach further comprising, simultaneous with step (e), mixing a tertiary DNA strand displacement primer with the amplification target circles (see col. 55 lines 1-10).

With regard to claim 20, Lizardi et al. teach further comprising detecting the tandem sequence DNA, wherein detection of tandem sequence DNA indicates that the corresponding messenger RNA molecule was present in the nucleic acid sample (see col. 52 lines 5-27).

With regard to claim 21, Lizardi et al. teach the tandem sequence DNA is detected while in association with the capture probes.

With regard to claim 22, Lizardi et al. teach the identity of the capture probe associated with a tandem sequence DNA indicates the identity of the corresponding messenger RNA molecule (see col. 52 lines 5-27).

With regard to claim 23, Lizardi et al. teach the tandem sequence DNA is detected at the site where the capture probe is located, and wherein the location of the capture probe indicates the identity of the corresponding messenger RNA molecule (see col. 52 lines 5-27).

With regard to claim 24, Lizardi et al. teach detection is mediated by detection probes or by a detection label incorporated in the tandem sequence DNA (see col. 48 lines 48-54).

With regard to claim 25, Lizardi et al. teach the detection label is a ligand (see col. 48 line 47). With regard to claim 26, Lizardi et al. teach the ligand is Brdu (see col. 48 lines 32-33).

With regard to claim 27, Lizardi et al. teach the ligand is Brdu, wherein the tandem sequence DNA is detected by associating an anti-Brdu antibody with the tandem sequence DNA and detecting the antiBrdu antibody (see col. 48 line 33).

With regard to claim 28, Lizardi et al. teach the anti-Brdu antibody comprises a label, wherein the anti-Brdu antibody is detected by detecting the label (see col. 48 lines 32-33).

With regard to claim 29, Lizardi et al. teach the label on the anti-Brdu antibody is a fluorophore (see col. 48 lines 32-37).

With regard to claim 31, Lizardi et al. teach further comprising mixing a set of detection probes with the tandem sequence DNA under conditions that promote hybridization between the tandem sequence DNA and the detection probes, and detecting a plurality of different sequences present in the tandem sequence DNA (see col. 61 lines 9-21).

With regard to claim 32, Lizardi et al. teach the tandem sequence DNA is collapsed using collapsing probes (see col. 52 lines 60-67).

Art Unit: 1637

With regard to claim 33, Lizardi et al. teach at least one of the collapsing probes is a collapsing detection probe (see col. 52 lines 22-23, 60).

With regard to claim 34, Lizardi et al. teach the tandem sequence DNA is collapsed by mixing the collapsing probes with the tandem sequence DNA, and incubating under conditions that promote hybridization between the collapsing probes and the tandem sequence DNA (see col. 52 lines 66-67, col. 55 line 14).

With regard to claim 35, Lizardi et al. teach further comprising, prior to or simultaneous with the mixing of the collapsing probes with the tandem sequence DNA, mixing detection probes with the tandem sequence DNA, and incubating under conditions that promote hybridization between the detection probes and the tandem sequence DNA (see col. 62 lines 66-67, col. 63 lines 1-10).

With regard to claim 36, Lizardi et al. teach the collapsing probes comprise ligands, haptens, or both coupled to or incorporated into oligonucleotides (see col. 63 lines 50-53, col. 24 lines 65-67).

With regard to claim 37, Lizardi et al. teach the RT primer comprises a capture tag (see col. 23 lines 50-67).

With regard to claims 38, 41 and 46, Lizardi et al. teach the biotin (see col. 23 lines 50-67).

With regard to claims 43 and 44, Lizardi et al. teach the association occurs between a protein and a nucleic acid (see col. 78 lines 31-32).

With regard to claim 61, Lizardi et al. teach the incorporation of biotinylated-ddNTP into the cDNA (see col. 43 lines 10-20).

Lizardi et al. do not teach the capture tag is a hapten a ligand a ligand binding molecule an antibody or an anti-antibody and the capture tag is not a nucleic acid.

Schweitzer et al. teach the capture tag is an antibiody.

With regard to claim 2, Schweitzer et al. teach the capture tag associates with the primer (see the abstract, where the primer is covalently attached to the antibody).

With regard to claims 39 and 40, Schweitzer et al. teach the DNA strands comprise capture tags (see the abstract, where the primer is covalently attached to the antibody).

With regard to claim 42, Schweitzer et al. teach the association is covalent (see the abstract, where the primer is covalently attached to the antibody).

With regard to claim 45, Schweitzer et al. teach the association occurs between two proteins (see p. 10116 col. 2 2<sup>nd</sup> full paragraph, where the analyte is PSA, a protein, and the tag bound to the primer is anti-PSA antibody, also a protein).

With regard to claims 54 and 57, Schweitzer et al. teach the rolling circle replication primers each comprise a capture tag (see the abstract, where the primer is covalently attached to the antibody).

With regard to claims 55 and 58, Schweitzer et al. teach association of the rolling circle replication primers with the DNA occurs via association of the capture tag added to the DNA and the capture tag in the rolling circle replication primers (see the abstract, where the primer is covalently attached to the antibody).

One of ordinary skill in the art at the time the invention was made would have been motivated to use the method of attaching an antibody capture tag to a primer, as taught by Schweitzer with the method of amplifying target nucleic acids as taught by Lizardi in order to have a versatile ultrasensitive method of antigen detection. Schweitzer teaches that the RCA reporter system can be adapted for the detection of protein antigens using an oligonulceotide primer that is covalently attached to an antibody. Schweitzer additionally, teaches using a single primer, RCA generates hundreds of tandemly linked copies of the circular template within a few minutes and the 5'end of the primer is attached to an antibody. The amplified DNA can be detected in a variety of ways including direct incorporation of hapten labeled or fluorescently labeled nucleotides. An ordinary practitioner would have been motivated to use the method of attaching an antibody capture tag to a primer, as taught by Schweitzer with the method of amplifying

target nucleic acids as taught by Lizardi in order to expand the genus of analytes which can be detected using RCA.

4. Claims 16 and 17 are rejected under 35 U.S.C. 103(a) as being unpatentable over Lizardi et al. (USPN 6,316,229 B1) and Schweitzer et al. (PNAS 2000) in view Lizardi (US 2003/0032024 A1).

The teachings and suggestions of Lizardi (229) and Schweitzer are described previously. Lizardi (229) and Schweitzer do not teach or suggest subprobes.

With regard to claims 16 and 17, Lizardi (024) teaches mixing one or more subprobes (gap oligonucleotides) with the cDNA strands wherein each half probe is designed to hybridize to a cDNA strand adjacent to where a capture probe hybridizes, ligating the subprobes and capture probes hybridized, and after ligation, incubating the capture probes at a temperature above the melting temperature of the capture probe but below the melting temperature of the ligated capture probe/subprobe (see paragraph 0195).

One of ordinary skill in the art at the time the invention was made one would have been motivated to apply Gap oligonucleotides as taught by Lizardi (024) with the method of amplifying target nucleic acids as taught by Lizardi (229) and Schweitzer in order to achieve more selective target discrimination. Lizardi (024) teaches using gap oligonucleotides enhance target dependency in LCR and this can be adapted for use in LM-RCA, and this method enhances target discrimination. It would have been prima facie obvious to use the gap oligonucleotide as taught by Lizardi (024) with the method of amplifying nucleic acids as taught by Lizardi (229) and Schweitzer in order to achieve the expected advantage of enhancing target nucleic acid discrimination.

3. Claim 30 is rejected under 35 U.S.C. 103(a) as being unpatentable over Lizardi et al. (USPN 6,316,229 B1) and Schweitzer et al. (PNAS 2000) in view of Waggoner et al. (USPN 6,008,373).

Art Unit: 1637

The teachings and suggestions of Lizardi (229) Schweitzer are described previously.

Lizardi (229) and Schweitzer do not teach or suggest phycoerythrin as a fluorophore.

Waggoner et al. teach using phycoerythrin as a fluorophore in the detection label on an antibody (see col. 21 line 64).

One of ordinary skill in the art at the time the invention was made one would have been motivated to use phycoerythrin as taught by Waggoner with the method of amplifying target nucleic acids as taught by Lizardi (229) and Schweitzer to achieve a detection signal that provides fluorescence that is relatively free of interference from other biological materials and provides a multicolor fluorescence emission using a single wavelength excitation. Waggoner teach phycoerythrin is advantageous because it is low molecular weight and provides a multicolor fluorescence emission using a single wavelength excitation (see col. 2 lines 28-30). It would have been prima facie obvious to use Phycoerythrin as taught by Waggoner with the method of amplifying nucleic acids as taught by Lizardi (229) and Schweitzer in order to achieve the expected advantage of a label that has a low molecular weight and provides a multicolor fluorescence emission from a single excitation wavelength.

4. Claims 48-52, and 69 are rejected under 35 U.S.C. 103(a) as being unpatentable over Lizardi et al. (USPN 6,316,229 B1) and Schweitzer et al. (PNAS 2000) in view of Cao et al. (US 2002/0120409 A1).

The teachings and suggestions of Lizardi (229) Schweitzer are described previously.

Lizardi (229) and Schweitzer do not teach or suggest fragmenting and labeling cDNA strands to form labeled fragmented cDNA.

Cao et al. teach fragmented cDNA in a method to amplify mRNA (see claim 1 page 8).

One of ordinary skill in the art at the time the invention was made one would have been motivated to use the method of fragmenting and labeling cDNA as taught by Cao et al. with the method of

amplifying target nucleic acids as taught by Lizardi (229) and Schweitzer to obtain labeled cDNA fragments that are used in assessing gene expression. It would have been prima facie obvious to use fragmenting and labeling cDNA as taught by Cao et al. with the method of amplifying target nucleic acids as taught by Lizardi (229) and Schweitzer in order to achieve the expected advantage of using the labeled cDNA fragments in gene expression arrays.

5. Claims 59-60 are rejected under 35 U.S.C. 103(a) as being unpatentable over Lizardi et al. (USPN 6,316,229 B1) and Schweitzer et al. (PNAS 2000) in view of Shoemaker et al. (USPN 6,713257 B2).

The teachings and suggestions of Lizardi (229) Schweitzer are described previously.

Lizardi (229) and Schweitzer do not teach or suggest teach a capture tag derived from allyl amine dUTP.

Shoemaker et al. teaches using an amino-allyl dUTP in labeling cDNA (see col. 34 line 8).

One of ordinary skill in the art at the time the invention was made one would have been motivated to use the method of labeling cDNA as taught by Shoemaker et al. with the method of amplifying target nucleic acids as taught by Lizardi (229) and Schweitzer to obtain labeled cDNA that are used in assessing gene expression. It would have been prima facie obvious to use the method of labeling cDNA as taught by Shoemaker et al. with the method of amplifying target nucleic acids as taught by Lizardi (229) and Schweitzer in order to achieve the expected advantage of incorporating a detectible fluorescent label into the cDNA of interest.

### Response to Arguments

6. Applicants' arguments filed March 22, 2006, have been fully considered but they are not persuasive.

With respect to Applicants' arguments regarding the combination of Lizardi (229) and Lizardi (024), these arguments are most in view of the new ground(s) of rejection.

With respect to Applicants' arguments regarding the limitation of RT primers, Applicants argue the 229 patent fails to disclose or refer specifically to RT primers. This is not persuasive because patent 229 expressly states that RNA can be used with the methodologies disclosed. Therefore a practioner of ordinary skill in the art recognizes the need for using RT primers when working with RNA. Additionally the 229 patent teaches cDNA which inherently requires the use of RT primers. Moreover primers are primers. RT primers (primers for producing cDNA) are not structurally different from DNA primers (primers for producing DNA), therefore because the combination of the 229 patent and 024 patent renders the instant invention obvious with respect to DNA primers then it necessarily renders the instant invention obvious with respect to RT primers.

## Summary

7. No claims were allowable.

## Correspondence

8. Any inquiry concerning this communication or earlier communications from the examiner should be directed to Heather G. Calamita whose telephone number is 571.272.2876 and whose e-mail address is heather.calamita@uspto.gov. However, the office cannot guarantee security through the e-mail system nor should official papers be transmitted through this route. The examiner can normally be reached on Monday through Thursday, 7:00 AM to 5:30 PM.

If attempts to reach the examiner are unsuccessful, the examiner's supervisor, Gary Benzion can be reached at 571.272.0782.

Papers related to this application may be faxed to Group 1637 via the PTO Fax Center using the fax number 571.273.8300.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to 571.272.0547.

Patent applicants with problems or questions regarding electronic images that can be viewed in the Patent Application Information Retrieval system (PAIR) can now contact the USPTO's Patent Electronic Business Center (Patent EBC) for assistance. Representatives are available to answer your questions daily from 6 am to midnight (EST). The toll free number is (866) 217-9197. When calling please have your application serial or patent number, the type of document you are having an image problem with, the number of pages and the specific nature of the problem. The Patent Electronic Business Center will notify applicants of the resolution of the problem within 5-7 business days. Applicants can also check PAIR to confirm that the problem has been corrected. The USPTO's Patent Electronic Business Center is a complete service center supporting all patent business on the Internet. The USPTO's PAIR system provides Internet-based access to patent application status and history information. It also enables applicants to view the scanned images of their own application file folder(s) as well as general patent information available to the public. For more information about the PAIR system, see http://pair-direct.uspto.gov.

For all other customer support, please call the USPTO Call Center (UCC) at 800-786-9199.

hgc

JEFFREY FREDMAN PRIMARY EXAMINER